



OIEP

RAW SEQUENCE LISTING
 PATENT APPLICATION: US/09/939,408A

DATE: 01/23/2002
 TIME: 17:03:26

Input Set : A:\500nscseq.txt
 Output Set: N:\CRF3\01232002\I939408A.raw

Does Not Contain
 Corrected Sequence

3 <110> APPLICANT: Yoshida, Roberta
 4 Kootstra, Anna
 7 <120> TITLE OF INVENTION: Phenylalanine Ammonia Lyase Polypeptide and
 8 Polynucleotide Sequences and Methods of Obtaining and
 9 Using Same
 11 <130> FILE REFERENCE: 29479/500NSCA
 13 <140> CURRENT APPLICATION NUMBER: US 09/939,408A
 17 <141> CURRENT FILING DATE: 2001-08-24
 19 <150> PRIOR APPLICATION NUMBER: US 09/624,693
 22 <151> PRIOR FILING DATE: 2000-07-24
 23 <150> PRIOR APPLICATION NUMBER: PCT/US01/23270
 26 <151> PRIOR FILING DATE: 2001-07-24
 29 <160> NUMBER OF SEQ ID NOS: 30
 31 <170> SOFTWARE: PatentIn Ver. 2.0

ERRORED SEQUENCES

1706 <210> SEQ ID NO: 21
 1707 <211> LENGTH: 726
 1708 <212> TYPE: PRT
 1709 <213> ORGANISM: Artificial Sequence
 1711 <220> FEATURE:
 1712 <221> NAME/KEY: SITE
 1713 <222> LOCATION: (12)..(719)
 1714 <223> OTHER INFORMATION: "Xaa"means any amino acid;"Xaa"means no consensus at that
 position
 1716 <223> OTHER INFORMATION: Description of Artificial Sequence: Consensus of
 1717 SEQ ID NOS: 13, 17, and 19
 1719 <400> SEQUENCE: 21
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 1722 1 5 10 15
 W--> 1724 Xaa Asn Gly Xaa His Ala Ala Xaa Xaa Ala Ser Xaa Xaa Xaa Xaa Xaa
 1725 20 25 30
 W--> 1727 Xaa Xaa Xaa Xaa Ala Xaa Ala Gly Ser Xaa Leu Pro Thr Thr Xaa Xaa
 1728 35 40 45
 W--> 1730 Thr Gln Leu Asp Ile Val Glu Xaa Xaa Leu Ala Asp Pro Xaa Thr Asp
 1731 50 55 60
 W--> 1733 Asp Xaa Xaa Glu Leu Asp Gly Tyr Ser Leu Thr Leu Gly Asp Val Val
 1734 65 70 75 80
 W--> 1736 Gly Ala Ala Arg Lys Gly Arg Xaa Val Arg Xaa Asp Ser Asp Glu
 1737 85 90 95
 W--> 1738

ile Arg Xaa Lys
 mal over

E--> 1739

100

105

110

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W--> 1741  Xaa Asn Ser Val Tyr Gly Val Thr Thr Gly Phe Gly Gly Ser Ala Asp
      1742      115      120      125
      1744  Thr Arg Thr Glu Asp Ala Ile Ser Leu Gln Lys Ala Leu Leu Glu His
      1745      130      135      140
W--> 1747  Gln Leu Cys Gly Val Leu Pro Thr Ser Xaa Asp Ser Phe Xaa Leu Gly
      1748      145      150      155      160
      1750  Arg Gly Leu Glu Asn Ser Leu Pro Leu Glu Val Val Arg Gly Ala Met
      1751      165      170      175
      1753  Thr Ile Arg Val Asn Ser Leu Thr Arg Gly His Ser Ala Val Arg Leu
      1754      180      185      190
      1756  Val Val Leu Glu Ala Leu Thr Asn Phe Leu Asn His Gly Ile Thr Pro
      1757      195      200      205
      1759  Ile Val Pro Leu Arg Gly Thr Ile Ser Ala Ser Gly Asp Leu Ser Pro
      1760      210      215      220
      1762  Leu Ser Tyr Ile Ala Ala Ala Ile Thr Gly His Pro Asp Ser Lys Val
      1763      225      230      235      240
W--> 1765  His Val Xaa His Glu Gly Xaa Glu Lys Ile Met Xaa Ala Arg Glu Ala
      1766      245      250      255
      1768  Ile Ala Leu Phe Gly Leu Glu Pro Val Val Leu Gly Pro Lys Glu Gly
      1769      260      265      270
      1771  Leu Gly Leu Val Asn Gly Thr Ala Val Ser Ala Ser Met Ala Thr Leu
      1772      275      280      285
      1774  Ala Leu His Asp Ala His Met Leu Ser Leu Leu Ser Gln Ala Leu Thr
      1775      290      295      300
      1777  Ala Leu Thr Val Glu Ala Met Val Gly His Ala Gly Ser Phe His Pro
      1778      305      310      315      320
      1780  Phe Leu His Asp Val Thr Arg Pro His Pro Thr Gln Ile Glu Val Ala
      1781      325      330      335
W--> 1783  Arg Asn Ile Arg Thr Leu Leu Glu Gly Ser Xaa Phe Ala Val His His
      1784      340      345      350
      1786  Glu Glu Glu Val Lys Val Lys Asp Asp Glu Gly Ile Leu Arg Gln Asp
      1787      355      360      365
      1789  Arg Tyr Pro Leu Arg Thr Ser Pro Gln Trp Leu Gly Pro Leu Val Ser
      1790      370      375      380
      1792  Asp Met Ile His Ala His Ala Val Leu Ser Leu Glu Ala Gly Gln Ser
      1793      385      390      395      400
W--> 1795  Thr Thr Asp Asn Pro Leu Ile Asp Val Glu Asn Lys Xaa Thr His His
      1796      405      410      415
      1798  Gly Gly Asn Phe Gln Ala Ser Ala Val Xaa Asn Thr Met Glu Lys Thr
      1799      420      425      430
      1801  Arg Leu Ala Leu Ala Leu Ile Gly Lys Leu Asn Phe Thr Gln Leu Thr
      1802      435      440      445
      1804  Glu Met Leu Asn Ala Gly Met Asn Arg Gly Leu Pro Ser Cys Leu Ala
      1805      450      455      460
      1807  Ala Glu Asp Pro Ser Leu Ser Tyr His Cys Lys Gly Leu Asp Ile Ala
      1808      465      470      475      480
      1810  Ala Ala Ala Tyr Thr Ser Glu Leu Gly His Leu Ala Asn Pro Val Thr
      1811      485      490      495
      1813  Thr His Val Gln Pro Ala Glu Met Gly Asn Gln Ala Val Asn Ser Leu

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1814          500          505          510
1816 Ala Leu Ile Ser Ala Arg Arg Thr Ala Glu Ala Asn Asp Val Leu Ser
1817          515          520          525
1817 Leu Leu Leu Ala Thr His Leu Tyr Cys Val Leu Gln Ala Val Asp Leu
1819          530          535          540
1820 Arg Ala Met Glu Phe Glu Phe Lys Lys Gln Phe Xaa Pro Xaa Xaa Xaa
W--> 1822          545          550          555          560
1823 Xaa Leu Leu Xaa Gln His Phe Gly Xaa Xaa Xaa Thr Xaa Xaa Xaa Xaa
W--> 1825          565          570          575          580
1826 Xaa Xaa Glu Leu Xaa Xaa Lys Val Xaa Lys Xaa Leu Xaa Lys Arg Leu
W--> 1828          585          590          595          600
1829 Glu Gln Thr Asn Ser Tyr Asp Leu Glu Pro Arg Trp His Asp Ala Phe
1831          605          610          615          620
1832 Ser Xaa Ala Thr Gly Thr Val Val Glu Xaa Leu Ser Ser Xaa Xaa Xaa
W--> 1834          625          630          635          640
1835 Xaa Xaa Val Ser Leu Ala Ala Val Asn Ala Trp Lys Val Ala Xaa Ala
W--> 1837          645          650          655          660
1838 Glu Lys Ala Ile Ser Leu Thr Arg Xaa Val Arg Xaa Xaa Phe Trp Xaa
W--> 1840          665          670          675          680
1841 Ala Pro Ser Ser Ser Ser Pro Ala Leu Xaa Tyr Leu Ser Pro Arg Thr
W--> 1843          685          690          695          700
1844 Arg Val Leu Tyr Ser Phe Val Arg Glu Glu Leu Gly Val Lys Ala Arg
1846          705          710          715          720
1847 Arg Gly Asp Val Phe Leu Gly Lys Gln Glu Val Thr Ile Gly Ser Asn
1849          725          730          735          740
1850 Val Ser Arg Ile Tyr Glu Ala Ile Lys Ser Gly Arg Ile Asn Xaa Val
W--> 1852          745          750          755          760
1853 Leu Val Lys Met Leu Ala
1855          765          770          775          780
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1921 <212> TYPE: DNA
1922 <213> ORGANISM: Rhodotorula graminis
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1925 <222> LOCATION: (6)
1926 <223> OTHER INFORMATION: Description of modified_base: m = a or c
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1928 <221> NAME/KEY: exon
1929 <222> LOCATION: (2008)..(2586)
1930 <223> OTHER INFORMATION:
W--> 1931 <220> FEATURE:
1932 <221> NAME/KEY: exon
1933 <222> LOCATION: (1822)..(1947)
1934 <223> OTHER INFORMATION:
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1936 <221> NAME/KEY: exon
1937 <222> LOCATION: (1587)..(1748)
1938 <223> OTHER INFORMATION:

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1942 <223> OTHER INFORMATION:
W--> 1943 <220> FEATURE:
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1946 <223> OTHER INFORMATION:
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1954 <223> OTHER INFORMATION:
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1956 <221> NAME/KEY: terminator
1957 <222> LOCATION: (2587)..(2589)
1958 <223> OTHER INFORMATION:
W--> 1959 <220> FEATURE:
1960 <221> NAME/KEY: Intron
1961 <222> LOCATION: (1948)..(2007)
1962 <223> OTHER INFORMATION:
W--> 1963 <220> FEATURE:
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1966 <223> OTHER INFORMATION:
W--> 1967 <220> FEATURE:
1968 <221> NAME/KEY: Intron
1969 <222> LOCATION: (1530)..(1586)
1970 <223> OTHER INFORMATION:
W--> 1971 <220> FEATURE:
1972 <221> NAME/KEY: Intron
1973 <222> LOCATION: (1296)..(1364)
1974 <223> OTHER INFORMATION:
W--> 1975 <220> FEATURE:
1976 <221> NAME/KEY: Intron
1977 <222> LOCATION: (881)..(960)
1978 <223> OTHER INFORMATION:
W--> 1979 <220> FEATURE:
1980 <221> NAME/KEY: Intron
1981 <222> LOCATION: (362)..(448)
1982 <223> OTHER INFORMATION:
W--> 1983 <400> SEQUENCE: 28
1984 atg gcm cct tcc ttg gac tcg ctc gcc acc acg ctc gcc aac ggc ttt 48
1985 Met Ala Pro Ser Leu Asp Ser Leu Ala Thr Thr Leu Ala Asn Gly Phe
E--> 1986
1 5 10
W--> 1987 Thr Asn Gly Ser His Ala Ala Pro Thr Lys Ser Ala Ala Gly Pro Thr
15 insert
15 insert return < acc aac ggc tcg c

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W--> 1988	20	25	30	144
1989	tgc gct ctc cgc cgc aag ccc ggc ctc gat ggc cac gcc gcg cac gag			
1990	Ser Ala Leu Arg Arg Thr Pro Gly Leu Asp Gly His Ala Ala His Gln			
W--> 1991	35	40	45	192
1992	tgc gag ctc gag atc gtg cag gag ctc ctc agc gac ccc acc gac gac			
1993	Ser Gln Leu Glu Ile Val Gln Glu Leu Ser Asp Pro Thr Asp Asp			
W--> 1994	50	55	60	240
1995	gtc gtc gag ctc agc ggc tac agc ctc acc gtc cgt gac gtt gtc ggc			
1996	Val Val Glu Leu Ser Gly Tyr Ser Leu Thr Val Arg Asp Val Val Gly			
W--> 1997	65	70	75	288
1998	gcc gcc cgc aag ggc cgc agg gtc cgc gtc cag aac gac gag atc			
1999	Ala Ala Arg Lys Gly Arg Arg Val Arg Val Gln Asn Asp Asp Gly Ile			
W--> 2000	85	90	95	336
2001	cgc gca cgc gtc gac aag agc gtc gac ttc ctc aag gcc cag ctt cag			
2002	Arg Ala Arg Val Asp Lys Ser Val Asp Phe Leu Lys Ala Gln Leu Gln			
W--> 2003	100	105	110	381
2004	aac tgc gtc tac gga gtc acc acg g tgcgttcga gacgagagc			
2005	Asn Ser Val Tyr Gly Val Thr Thr			
W--> 2006	115	120		441
2007	ggaaatctcg ggaatgcgcga gcgtgaacg ctgacactcg cttggacggc tgcgcggctc			489
2008	ttgcagg gt ttc ggt ggc tgc gcc gac acg agg acc gag gat gca gtc			
2009	Gly Phe Gly Gly Ser Ala Asp Thr Arg Thr Glu Asp Ala Val			
W--> 2010	125	130		537
2011	agc ctc cag aag gcg ctc atc gag cac cag ctc tgc gcc gtg acg ccg			
2012	Ser Leu Gln Lys Ala Leu Ile Glu His Gln Leu Cys Gly Val Thr Pro			
W--> 2013	135	140	145	585
2014	acg tcc gtc tgc tcc ttc agc gtc gga cgc gcc ctc gag aac acg ctt			
2015	Thr Ser Val Ser Ser Phe Ser Val Gly Arg Gly Leu Glu Asn Thr Leu			
W--> 2016	155	160	165	633
2017	cgc ctc gag gtc gtc cgc gcc gcc atg gtc atc cgc gtc aac tgc ctc			
2018	Pro Leu Glu Val Val Arg Gly Ala Met Val Ile Arg Val Asn Ser Leu			
W--> 2019	170	175	180	681
2020	acg cgt gcc cac tgc gcc gtc cgc ctc gtc gtc ctt gag gcg ctc acc			
2021	Thr Arg Gly His Ser Ala Val Arg Leu Val Val Leu Glu Ala Leu Thr			
W--> 2022	185	190	195	729
2023	aac ttc ttg aac cac cgc atc acg ccc atc gtc ccc ctc cgc gcc tcc			
2024	Asn Phe Leu Asn His Arg Ile Thr Pro Ile Val Pro Leu Arg Gly Ser			
W--> 2025	200	205	210	777
2026	atc tgc gcg tgc gcc gac ctc agc ccg ctc tgc tac atc gcc gcc gcc			
2027	Ile Ser Ala Ser Gly Asp Leu Ser Pro Leu Ser Tyr Ile Ala Gly Ala			
W--> 2028	215	220	225	825
2029	atc acc ggt cac ccc gac gtc aag gtt cac gtt ttg cac gag gga acc			
2030	Ile Thr Gly His Pro Asp Val Lys Val His Val Leu His Glu Gly Thr			
W--> 2031	235	240	245	873
2032	gag aag atc atg ttt gcg cgc gag gcc atc tgc ctc ttt ggt ctc gag			
2033	Glu Lys Ile Met Phe Ala Arg Glu Ala Ile Ser Leu Phe Gly Leu Glu			
W--> 2034	250	255	260	930
2035	gca gtc g gtaactgcgc agtctcgact gcagtgcgct gtctgcagagt ctccccagtt			
2036	Ala Val			

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W--> 2038      gctgactgcc ctttgttcac gcgattgcag tc ctc ggc cgc aag gag ggt ctc 983
2039                      Val Leu Gly Pro Lys Glu Gly Leu
W--> 2040                      270
2041      ggt ctg gtc aac gga acg gcc gtc tcc gcc tcg atg gcg acc ctc agt 1031
2042      Gly Leu Val Asn Gly Thr Ala Val Ser Ala Ser Met Ala Thr Leu Ser
2043                      275
2044      ctg cac gac tcg cac atg ctc tcg ctc tcg cag gcc ttg acg gct 1079
2045      Leu His Asp Ser His Met Leu Ser Leu Leu Ser Gln Ala Leu Thr Ala
2046                      290
2047      ctc acg gtg gag gcc atg gtc ggc cag cag ggc tcg ttc gcg cgc ttc 1127
2048      Leu Thr Val Glu Ala Met Val Gly Gln Gln Gly Ser Phe Ala Pro Phe
2049                      305
2050      atc cac gac gtc tgc cgc cgc ccc gcc cag gtc gag gtc gcg cgc 1175
2051      Ile His Asp Val Cys Arg Pro His Pro Gly Gln Val Glu Val Ala Arg
2052                      325
2053      aac atc cgc acg ctc ctt tcc ggc tcg tcg ttt gcc gtt gag cac gag 1223
2054      Asn Ile Arg Thr Leu Leu Ser Gly Ser Phe Ala Val Glu His Glu
2055                      340
2056      gag gag gtc aag gtc aag gac gag gag ggc att ctt cgc cag gac cgc 1271
2057      Glu Glu Val Lys Val Lys Asp Asp Glu Gly Ile Leu Arg Gln Asp Arg
2058                      355
2059      tac cgc ctc cgc acg tcg cct cag gttcgtcccc tctctctccc cttccctcgc 1325
2060      Tyr Pro Leu Arg Thr Ser Pro Gln
2061                      375
2062      tccgaccggc gcgtcgagac ttacgttttg cgtatccag ttc ctc ggc cgc ctc 1379
2063                      380
2064      gtg gag gac atg atg cac gcc tac tcg act ctc tcg ctc gag aac aac 1427
2065      Val Glu Asp Met Met His Ala Tyr Ser Thr Leu Ser Leu Glu Asn Asn
2066                      385
2067      acg acg acc gac aac cgc ctc ctc gac gtc gag aac aag cag acc gcg 1475
2068      Thr Thr Thr Asp Asn Pro Leu Leu Asp Val Glu Asn Lys Gln Thr Ala
2069                      400
2070      cac ggc ggc aac ttc cag gcg tcg gct gtc tcg att tcg atg gag aag 1523
2071      His Gly Gly Asn Phe Gln Ala Ser Ala Val Ser Ile Ser Met Glu Lys
2072                      415
2073      acc agg tgcgtctctc gctgccttcg tactccgatac ttgtgctgaa tgtttctctc 1579
2074                      420
2075      ctgcagg ctc gca ctc gcc ctc atc ggc aag ctc aac ttc acg cag tgc 1628
2076      Leu Ala Leu Ala Leu Ile Gly Lys Leu Asn Phe Thr Gln Cys
2077                      435
2078      acc gag ttg ctc aac gct gcc atg aac cgc ggc ctg cct tcg tgc ctc 1676
2079      Thr Glu Leu Leu Asn Ala Ala Met Asn Arg Gly Leu Pro Ser Cys Leu
2080                      450
2081      gct gcc gag gac cgc tcg ctc aac tat cac ggc aag ggc ttg gac att 1724
2082      Ala Ala Glu Asp Pro Ser Leu Asn Tyr His Gly Lys Gly Leu Asp Ile
2083                      465
2084      cac atc gct gct tac gct tcg gag gtgagcgcgc gacgttctcc gccctgcctc 1778
2085                      470
2086

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2137	aac cac gtc ctc gtc aag atg ctc gcg taaggcccg gcaagcctcg	2606
2138	Asn His Val Leu Val Lys Met Leu Ala	
2139	715 720	
2140	cctagacgcc cgctcaccc caagaccagc ttttcgacgt cgtgtcgtgc caagaacgga	2666
2141	ctttctcca tacacatgct gtcttactct ctgcgcgtca tcaegtctct cagttctttc	2726
2142	gtatcccgcg tctct	2741

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L:61 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2
L:79 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3
L:197 M:283 W: Missing Blank Line separator, <220> field identifier
L:252 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12
L:416 M:283 W: Missing Blank Line separator, <400> field identifier
L:444 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:13
L:1622 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:1624 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:1626 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
L:1628 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:20
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L:1730 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21

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 L:1736 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
 L:1738 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21
 L:1739 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:21
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 L:1906 M:220 C: Keyword misspelled or invalid format, <213> ORGANISM for SEQ ID#:26
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 L:1909 M:283 W: Missing Blank Line separator, <400> field identifier
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 L:1971 M:283 W: Missing Blank Line separator, <220> field identifier
 L:1975 M:283 W: Missing Blank Line separator, <220> field identifier
 L:1979 M:283 W: Missing Blank Line separator, <400> field identifier
 L:1983 M:283 W: Missing Blank Line separator, in Coding Region, SEQ ID:28
 L:1986 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:1986 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:6
 L:1987 M:361 W: Invalid Split Codon, Sequence data for SEQ ID#: 28
 L:1988 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:1991 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:1994 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:1997 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2000 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2003 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2006 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2010 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2013 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2016 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2019 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2022 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2025 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2028 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2031 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2034 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2038 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2040 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:28
 L:2148 M:283 W: Missing Blank Line separator, <220> field identifier

VERIFICATION SUMMARY
PATENT APPLICATION: US/09/939,408ADATE: 01/23/2002
TIME: 17:03:27Input Set : A:\500nscseq.txt
Output Set : N:\CRF3\01232002\I939408A.raw

L:2151 M:283 W: Missing Blank Line separator, <220> field identifier
L:2155 M:283 W: Missing Blank Line separator, <400> field identifier
L:2296 M:283 W: Missing Blank Line separator, <220> field identifier
L:2299 M:283 W: Missing Blank Line separator, <220> field identifier
L:2303 M:283 W: Missing Blank Line separator, <220> field identifier
L:2307 M:283 W: Missing Blank Line separator, <220> field identifier
L:2311 M:283 W: Missing Blank Line separator, <220> field identifier
L:2315 M:283 W: Missing Blank Line separator, <220> field identifier
L:2319 M:283 W: Missing Blank Line separator, <220> field identifier
L:2323 M:283 W: Missing Blank Line separator, <220> field identifier
L:2327 M:283 W: Missing Blank Line separator, <220> field identifier
L:2331 M:283 W: Missing Blank Line separator, <220> field identifier
L:2335 M:283 W: Missing Blank Line separator, <220> field identifier
L:2339 M:283 W: Missing Blank Line separator, <220> field identifier
L:2343 M:283 W: Missing Blank Line separator, <220> field identifier
L:2347 M:283 W: Missing Blank Line separator, <220> field identifier
L:2351 M:283 W: Missing Blank Line separator, <220> field identifier
L:2355 M:283 W: Missing Blank Line separator, <220> field identifier
L:2359 M:283 W: Missing Blank Line separator, <220> field identifier
L:2363 M:283 W: Missing Blank Line separator, <220> field identifier
L:2367 M:283 W: Missing Blank Line separator, <220> field identifier
L:2371 M:283 W: Missing Blank Line separator, <220> field identifier
L:2375 M:283 W: Missing Blank Line separator, <220> field identifier
L:2379 M:283 W: Missing Blank Line separator, <220> field identifier
L:2383 M:283 W: Missing Blank Line separator, <220> field identifier
L:2387 M:283 W: Missing Blank Line separator, <220> field identifier
L:2391 M:283 W: Missing Blank Line separator, <220> field identifier